

Supporting Information

Survival of phenotypic information during cellular growth transitions

J. Christian J. Ray^{*}

^{*}Corresponding author.

Address:

Center for Computational Biology
Department of Molecular Biosciences
University of Kansas
2030 Becker Dr.
Lawrence, KS 66047

Email: jjray@ku.edu

Tel: 785-864-1506

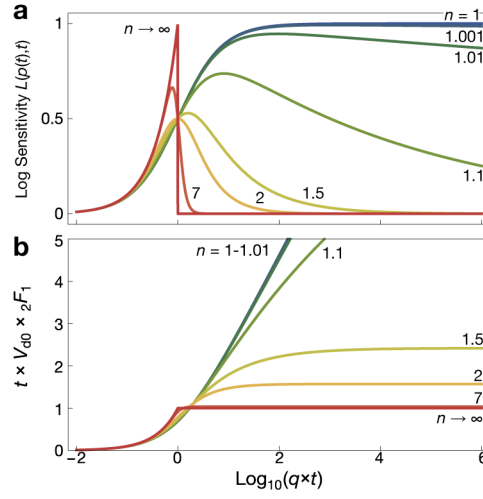


Figure S1. Effects of changing the steepness parameter in the growth arrest rate, n , on protein dynamics with negligible growth-independent protein loss ($\delta = 0$). **a.** Logarithmic change in protein concentration over time for various values of steepness parameter n . **b.** Behavior of the generalized growth-mediated dilution term, $tV_{d0} {}_2F_1\left(1, \frac{1}{n}; 1 + \frac{1}{n}; -(qt)^n\right)$, for various values of n .

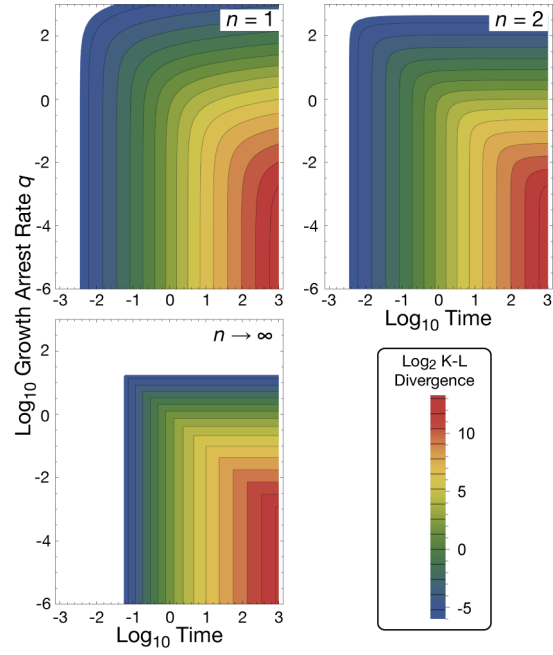


Figure S2. Kullback-Leiber divergence, in units of bits, for $n = 1, 2$, and $n \rightarrow \infty$. White regions, $D < 2^{-6} = 1/64$. Computed from Eqn. (9) for a gamma distribution. For sufficiently sharp and fast growth arrest transitions, the divergence is negligible, or stops increasing at low values, as protein turnover is halted.